

#3

0450

OIPF

RAW SEQUENCE LISTING

DATE: 02/08/2001

PATENT APPLICATION: US/09/769,878

TIME: 12:15:20

Input Set : A:\2976-B seq.txt

Output Set: N:\CRF3\02082001\I769878.raw

ENTERED

3 <110> APPLICANT: Sims, John
 5 <120> TITLE OF INVENTION: FIL-1 THETA DNAs AND POLYPEPTIDES
 7 <130> FILE REFERENCE: 2976-B
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/769,878
 10 <141> CURRENT FILING DATE: 2001-01-25
 12 <150> PRIOR APPLICATION NUMBER: US 60/195,962
 13 <151> PRIOR FILING DATE: 2000-04-11
 15 <150> PRIOR APPLICATION NUMBER: US 60/178,389
 16 <151> PRIOR FILING DATE: 2000-01-27
 18 <160> NUMBER OF SEQ ID NOS: 21
 20 <170> SOFTWARE: PatentIn version 3.0
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 339
 24 <212> TYPE: DNA
 25 <213> ORGANISM: Homo sapien
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 30 atccaggag ggagccgctg cctggcatgt gtggagacag aagaggggccc ttccctacag 120
 32 ctggaggatg tgaacattga ggaactgtac aaaggtgggtg aagagggccac acgcttcacc 180
 34 ttcttccaga gcagctcagg ctccgccttc aggcttgagg ctgtgcctg gcctggctg 240
 36 ttctgtgtg gcccggcaga gcccagcag ccagtacagc tcaccaagga gagtgagccc 300
 38 toagcccgtg ccaagtttta ctttgaacag agctggtag 339
 41 <210> SEQ ID NO: 2
 42 <211> LENGTH: 112
 43 <212> TYPE: PRT
 44 <213> ORGANISM: Homo sapiens
 46 <400> SEQUENCE: 2
 48 Lys Ile Cys Ile Leu Pro Asn Arg Gly Leu Ala Arg Thr Lys Val Pro
 49 1 5 10 15
 51 Ile Phe Leu Gly Ile Gln Gly Gly Ser Arg Cys Leu Ala Cys Val Glu
 52 20 25 30
 54 Thr Glu Glu Gly Pro Ser Leu Gln Leu Glu Asp Val Asn Ile Glu Glu
 55 35 40 45
 57 Leu Tyr Lys Gly Gly Glu Glu Ala Thr Arg Phe Thr Phe Phe Gln Ser
 58 50 55 60
 60 Ser Ser Gly Ser Ala Phe Arg Leu Glu Ala Ala Ala Trp Pro Gly Trp
 61 65 70 75 80
 63 Phe Leu Cys Gly Pro Ala Glu Pro Gln Gln Pro Val Gln Leu Thr Lys
 64 85 90 95
 66 Glu Ser Glu Pro Ser Ala Arg Thr Lys Phe Tyr Phe Glu Gln Ser Trp
 67 100 105 110
 69 <210> SEQ ID NO: 3
 70 <211> LENGTH: 459
 71 <212> TYPE: DNA
 72 <213> ORGANISM: Homo sapien
 74 <220> FEATURE:
 75 <221> NAME/KEY: CDS

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78 <400> SEQUENCE: 3
79 atg tgt tcc ctc ccc atg gca aga tac tac ata att aaa tat gca gac      48
80 Met Cys Ser Leu Pro Met Ala Arg Tyr Tyr Ile Ile Lys Tyr Ala Asp
81 1          5          10          15
83 cag aag gct cta tac aca aga gat ggc cag ctg ctg gtg gga gat cct      96
84 Gln Lys Ala Leu Tyr Thr Arg Asp Gly Gln Leu Leu Val Gly Asp Pro
85          20          25          30
87 gtt gca gac aac tgc tgt gca gag aag atc tgc aca ctt cct aac aga      144
88 Val Ala Asp Asn Cys Cys Ala Glu Lys Ile Cys Thr Leu Pro Asn Arg
89          35          40          45
91 ggc ttg gac cgc acc aag gtc ccc att ttc ctg ggg atc cag gga ggg      192
92 Gly Leu Asp Arg Thr Lys Val Pro Ile Phe Leu Gly Ile Gln Gly Gly
93          50          55          60
95 agc cgc tgc ctg gca tgt gtg gag aca gaa gag ggg cct tcc cta cag      240
96 Ser Arg Cys Leu Ala Cys Val Glu Thr Glu Glu Gly Pro Ser Leu Gln
97 65          70          75          80
99 ctg gag gat gtg aac att gag gaa ctg tac aaa ggt ggt gaa gag gcc      288
100 Leu Glu Asp Val Asn Ile Glu Glu Leu Tyr Lys Gly Gly Glu Glu Ala
101          85          90          95
103 aca cgc ttc acc ttc ttc cag agc agc tca ggc tcc gcc ttc agg ctt      336
104 Thr Arg Phe Thr Phe Phe Gln Ser Ser Ser Gly Ser Ala Phe Arg Leu
105          100          105          110
107 gag gcc gct gcc tgg cct ggc tgg ttc ctg tgt ggc ccg gca gag ccc      384
108 Glu Ala Ala Ala Trp Pro Gly Trp Phe Leu Cys Gly Pro Ala Glu Pro
109          115          120          125
111 cag cag cca gta cag ctc act aag gag agt gag ccc tca gcc cgt acc      432
112 Gln Gln Pro Val Gln Leu Thr Lys Glu Ser Glu Pro Ser Ala Arg Thr
113          130          135          140
115 aag ttt tac ttt gaa cag agc tgg tag      459
116 Lys Phe Tyr Phe Glu Gln Ser Trp
117 145          150
120 <210> SEQ ID NO: 4
121 <211> LENGTH: 152
122 <212> TYPE: PRT
123 <213> ORGANISM: Homo sapien
125 <400> SEQUENCE: 4
127 Met Cys Ser Leu Pro Met Ala Arg Tyr Tyr Ile Ile Lys Tyr Ala Asp
128 1          5          10          15
131 Gln Lys Ala Leu Tyr Thr Arg Asp Gly Gln Leu Leu Val Gly Asp Pro
132          20          25          30
135 Val Ala Asp Asn Cys Cys Ala Glu Lys Ile Cys Thr Leu Pro Asn Arg
136          35          40          45
139 Gly Leu Asp Arg Thr Lys Val Pro Ile Phe Leu Gly Ile Gln Gly Gly
140          50          55          60
143 Ser Arg Cys Leu Ala Cys Val Glu Thr Glu Glu Gly Pro Ser Leu Gln
144 65          70          75          80
147 Leu Glu Asp Val Asn Ile Glu Glu Leu Tyr Lys Gly Gly Glu Glu Ala
148          85          90          95

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151 Thr Arg Phe Thr Phe Phe Gln Ser Ser Ser Gly Ser Ala Phe Arg Leu
152                100                105                110
155 Glu Ala Ala Ala Trp Pro Gly Trp Phe Leu Cys Gly Pro Ala Glu Pro
156                115                120                125
159 Gln Gln Pro Val Gln Leu Thr Lys Glu Ser Glu Pro Ser Ala Arg Thr
160                130                135                140
163 Lys Phe Tyr Phe Glu Gln Ser Trp
164 145                150
167 <210> SEQ ID NO: 5
168 <211> LENGTH: 21
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial
172 <220> FEATURE:
173 <223> OTHER INFORMATION: DNA primer
175 <400> SEQUENCE: 5
176 agaagatctg catacttct a                                21
179 <210> SEQ ID NO: 6
180 <211> LENGTH: 19
181 <212> TYPE: DNA
182 <213> ORGANISM: Artificial
184 <220> FEATURE:
185 <223> OTHER INFORMATION: DNA primer
187 <400> SEQUENCE: 6
188 tgagcaggat gagcttggt                                19
191 <210> SEQ ID NO: 7
192 <211> LENGTH: 27
193 <212> TYPE: DNA
194 <213> ORGANISM: Artificial
196 <220> FEATURE:
197 <223> OTHER INFORMATION: DNA primer
199 <400> SEQUENCE: 7
200 ccatacctaatacgcactcact atagggc                                27
203 <210> SEQ ID NO: 8
204 <211> LENGTH: 18
205 <212> TYPE: DNA
206 <213> ORGANISM: Artificial
208 <220> FEATURE:
209 <223> OTHER INFORMATION: DNA primer
211 <400> SEQUENCE: 8
212 ctccagctgt agggaagg                                18
215 <210> SEQ ID NO: 9
216 <211> LENGTH: 23
217 <212> TYPE: DNA
218 <213> ORGANISM: Artificial
220 <220> FEATURE:
221 <223> OTHER INFORMATION: DNA primer
223 <400> SEQUENCE: 9
224 actcactata gggctcgagc ggc                                23
227 <210> SEQ ID NO: 10

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RAW SEQUENCE LISTING

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Input Set : A:\2976-B seq.txt

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230 <213> ORGANISM: Artificial
232 <220> FEATURE:
233 <223> OTHER INFORMATION: DNA primer
235 <400> SEQUENCE: 10
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239 <210> SEQ ID NO: 11
240 <211> LENGTH: 20
241 <212> TYPE: DNA
242 <213> ORGANISM: Artificial
244 <220> FEATURE:
245 <223> OTHER INFORMATION: DNA primer
247 <400> SEQUENCE: 11
248 caaggctaaa acgcagtttc 20
251 <210> SEQ ID NO: 12
252 <211> LENGTH: 20
253 <212> TYPE: DNA
254 <213> ORGANISM: Artificial
256 <220> FEATURE:
257 <223> OTHER INFORMATION: DNA primer
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264 <211> LENGTH: 22
265 <212> TYPE: DNA
266 <213> ORGANISM: Artificial
268 <220> FEATURE:
269 <223> OTHER INFORMATION: DNA primer
271 <400> SEQUENCE: 13
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276 <211> LENGTH: 538
277 <212> TYPE: DNA
278 <213> ORGANISM: Mus musculus
280 <220> FEATURE:
281 <221> NAME/KEY: CDS
282 <222> LOCATION: (29)..(487)
284 <400> SEQUENCE: 14
285 tgaagaccag acactoccaa ctgcagga atg tgc tcc ctt ccc atg gca aga 52
286 Met Cys Ser Leu Pro Met Ala Arg
287 1 5
289 tac tac ata atc aag gat gca cat caa aag gct ttg tac aca cgg aat 100
290 Tyr Tyr Ile Ile Lys Asp Ala His Gln Lys Ala Leu Tyr Thr Arg Asn
291 10 15 20
293 ggc cag ctc ctg ctg gga gac cct gat tca gac aat tat agt cca gag 148
294 Gly Gln Leu Leu Leu Gly Asp Pro Asp Ser Asp Asn Tyr Ser Pro Glu
295 25 30 35 40
297 aag gtc tgt atc ctt cct aac cga ggc cta gac cgc tcc aag gtc ccc 196

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298 Lys Val Cys Ile Leu Pro Asn Arg Gly Leu Asp Arg Ser Lys Val Pro
299          45          50          55
301 atc ttc ctg ggg atg cag gga gga agt tgc tgc ctg gcg tgt gta aag      244
302 Ile Phe Leu Gly Met Gln Gly Gly Ser Cys Cys Leu Ala Cys Val Lys
303          60          65          70
305 aca aga gag gga cct ctc ctg cag ctg gag gat gtg aac atc gag gac      292
306 Thr Arg Glu Gly Pro Leu Leu Gln Leu Glu Asp Val Asn Ile Glu Asp
307          75          80          85
309 cta tac aag gga ggt gaa caa acc acc cgt ttc acc ttt ttc cag aga      340
310 Leu Tyr Lys Gly Gly Glu Gln Thr Thr Arg Phe Thr Phe Phe Gln Arg
311          90          95          100
313 agc ttg gga tct gcc ttc agg ctt gag gct gct gcc tgc cct ggc tgg      388
314 Ser Leu Gly Ser Ala Phe Arg Leu Glu Ala Ala Ala Cys Pro Gly Trp
315 105          110          115          120
317 ttt ctc tgt ggc cca gct gag ccc cag cag cca gtg cag ctc acc aaa      436
318 Phe Leu Cys Gly Pro Ala Glu Pro Gln Gln Pro Val Gln Leu Thr Lys
319          125          130          135
321 gag agt gaa ccc tcc acc cat act gaa ttc tac ttt gag atg agt cgg      484
322 Glu Ser Glu Pro Ser Thr His Thr Glu Phe Tyr Phe Glu Met Ser Arg
323          140          145          150
325 taa ggagacataa ggctggggcc tcgtctagtgc cccccagtct gagatcttct t      538
328 <210> SEQ ID NO: 15
329 <211> LENGTH: 152
330 <212> TYPE: PRT
331 <213> ORGANISM: Mus musculus
333 <400> SEQUENCE: 15
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336 1          5          10          15
339 Gln Lys Ala Leu Tyr Thr Arg Asn Gly Gln Leu Leu Leu Gly Asp Pro
340          20          25          30
343 Asp Ser Asp Asn Tyr Ser Pro Glu Lys Val Cys Ile Leu Pro Asn Arg
344          35          40          45
347 Gly Leu Asp Arg Ser Lys Val Pro Ile Phe Leu Gly Met Gln Gly Gly
348          50          55          60
351 Ser Cys Cys Leu Ala Cys Val Lys Thr Arg Glu Gly Pro Leu Leu Gln
352 65          70          75          80
355 Leu Glu Asp Val Asn Ile Glu Asp Leu Tyr Lys Gly Gly Glu Gln Thr
356          85          90          95
359 Thr Arg Phe Thr Phe Phe Gln Arg Ser Leu Gly Ser Ala Phe Arg Leu
360          100          105          110
363 Glu Ala Ala Ala Cys Pro Gly Trp Phe Leu Cys Gly Pro Ala Glu Pro
364          115          120          125
367 Gln Gln Pro Val Gln Leu Thr Lys Glu Ser Glu Pro Ser Thr His Thr
368          130          135          140
371 Glu Phe Tyr Phe Glu Met Ser Arg
372 145          150
375 <210> SEQ ID NO: 16
376 <211> LENGTH: 30
377 <212> TYPE: DNA

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/769,878

DATE: 02/08/2001

TIME: 12:15:21

Input Set : A:\2976-B seq.txt

Output Set: N:\CRF3\02082001\I769878.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number